ATG MET CCC316 AGG TTT CCC GAC TGG AAA GCG GTC AGT GAG CGC AAC ACA ATT AAT GTG AGT TAG CTC ACT CAT GGA TIT CAC ACA GGA AAC AGC TAT GAC CAT GAT TAC GTC CAA GCT CGA AAT TAA CCC TCA 190 CTT TCG CCC GGA AGG GGA CCG ACC GAG CCC AGC GAT TCA TGG AGC GAG GAA AGC GGG AAG AGC GCC CAA TAC CCA AGC CGC CTC TCG CCG CGT TGT GCG ATT CAT TAA TAC AGC TGC CAC GAC 127 TTA ATC CCC CTG ATT CCT TGT GGG ATA ACC CGT ATT CCC CCC CTT AGA GTG AAT TTG AAA ACC 1 GAT TIT TIG TGA TAT CIT CIG GGG GGG GGG GAA CCT AIT GTA TAA ACG CCA ACC AAC CGG TTT GGG TAC CTG GCC ATT TTA CTT GGC CCA TTT TGG TAA AAT GTT CCT TTC CCT GCG 505 CTA AAG GGA ACA AAA ACT GGA GCT CCA CCG CGG TGG CGG CCG CTC TAG AAC TAG TGG ATC 379 TAG GCA CCC CAG GCT TTA CAC TTT ATG CTT CCG GCT CGT ATG TTG TGT GGA ATT GTG AGC ø 568 CGG GCT CCA GGA ATT CGC CAC GAC CGG GAG GAC CGA GGA GGC GCC AGA CTA CGG GCG TAA CAA 64 CCC 253

TTC Trp Pro Gly Tyr Asp Leu Asn Leu Phe \mathtt{TTA} TTG AAT GTG ATT ATG GAT GAT TGG CCA GGG TAT GAC Asp MET Asp Val Ile GIC Ala Thr Arg Ser Pro Gly Val CCT GGC 629 GCG ACC CGC AGC

FIG. 1A

- Ile ATTIle ATC G1yGGT His CAT Pro CCT Ile ATC Leu CTC Val GTC TyrTAT Glu GAG Leu TTG Asp GAC GlyGGA TyrTAT TyrTAT His CAC Gln CAG Pro CCA TyrTAC Thr 692
- Val MET ATG ATC Ile Asp GAC Ser AGT Tyr TAT GlyGGA Lys Asp Ile ATA GAC AAA MET ATG ATT Asp GAT Lys AAG Ala CCC Leu CTG Arg CGG Glu GAG Ile ATT Arg AGA GAC 755
- Asn AAC Lys AAG Len CTT CAC His Glu GAA GTA Val CTC Leu Asp GAT Ala GCT Cys TGT Phe TTC LysAAA TAC $\frac{\text{GGG}}{\text{G1y}}$ G1y999 LysAAA Leu CTT Val GTG CysTGT CTG Len 818
- Arg AGG Tyr TAC Ser AGT Lys AAA Leu CTA AGA Arg ATC Ile TTC Asp Phe GAT GTT Val Lys AAG ATG MET TCA Ser <u>GTC</u> Val Phe TTT Arg 990 Asp GAT Ser TCA Asn AAT CGA Ser AGC 881
- GGA GlyAla GCTLeu CTG Thr ACG TCA Ser CTTLeu Asp GAT GGT G1y<u>GGC</u> G1y G1yGGA Ile ATC ATA Ile Gln CAG ATG MET GAG Glu GGT Gly ATG MET TCC Ser CAG Gln Asp GAC Asn AAT 944
- AGC Ser Leu CTCLeu CTA Ala GCA LysAAA ATG MET ACC Thr AGG Arg <u>666</u> G1y ACT GGA G1yGTC Val GTT Val GAT Asp GAG Glu GTT Val Ile ATT CTC Len TTT Phe AAT Asn AAG 1007
- TCC Ser ACA Arg AGA AAG Lys GTG Val Leu TTG TTGLeu Ser AGT Ala 205 GTA Val AAG Lys Ile ATT ATG MET AAC Asn 222 Pro Lys AAG $\frac{\text{TAC}}{\text{TYr}}$ Lys AAA GAG Glu ATA AAT 1070
- GGA GlyVal GTG Val GTG Phe TTT Leu TTA His CAC Pro CCA Ile ATT Glu GAG Phe $_{
 m LLL}$ GGA G1yAla GCT TyrTAT GAC Phe Arg Pro Asp CCT AGA TTT GlyCGC Asp GAC Ser AGT AGA

FIG. 1B

- 1196 TAT GCC TTA GAT TAC AAT GAA TAC TTC AGA GAT CTG AAT CAC ATA TGC GTC ATC AAT GAG CAC TYR Ala Leu Asp Tyr Asn Glu Tyr Phe Arg Asp Leu Asn His Ile Cys Val Ile Asn Glu His
- GGG TAA AGG AAA ATA TCG AGT CTT AAA GAC ATG AAT TCT CAC CAC TAA AGG CCC CAG ATA GGA Gly STP 1259
- 1322 TCA TTT TTA CGC CTG TCT TGG GGA GCC AGT TGC AAG TTG GGC CCC CCC GGA TCT TCA TCA GGA

1385 GG

1	MATRSPGVVISDDEPGYDLDLFCIPNHYAEDLERVFIPHGLIMDRTERLARDVMKE	56
1		56
57	MGGHHIVALCVLKGGYKFFADLLDYIKALNRNSDRSIPMTVDFIRLKSYCNDQSTG	112
57	: . :.	112
113	DIKVIGGDDLSTLTGKNVLIVEDIIDTGKTMQTLLSLVRQYNPKMVKVASLLVKRT :::: : : : : : : : : : : :	168
113	EMQIIGGGDLSTLAGKNFLIVEDVVGTGRTMKALLSNIEKYKPNMIKVASLLVKRT	168
169	PRSVGYKPDFVGFEIPDKFVVGYALDYNEYFRDLNHVCVISETGKAKYKA	218
169	. :: :. .	218

FIG.2

Majority HPRT2b.pep HPRThu HPRTcl HPRTplas HPRTtrypan	Majority	HPRT2b.pep HPRThu HPRTcl HPRTplas HPRTtrypan	Majority	HPRT2b.pep HPRThu HPRTcl HPRTplas HPRTtyrpan
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FIG.3A

Majority HPRT2b.pep HPRThu HPRTplas HPRTplas	Majority HPRT2b.pep HPRThu HPRTcl HPRTplas	Majority HPRT2b.pep HPRThu HPRTplas HPRTtrypan
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FIG.3B

Majority HPRT2b.pep HPRThu HPRTcl	HPRTplas HPRTtrypan	Majority	HPRT2.pep HPRThu	HPRTcl HPRTplas	HPRTtrypan	Majority	HPRT2b.pep HPRThu	HPRTcl HPRTplas HPRTtrypan
R S V G F K P D F V G F E I P D K F V V G Y A L D Y N E Y F 190 R S D G F R P D Y A G F E I P H L F V V G Y A L D Y N E Y F R S V G Y K P D F V G F E I P D K F V V G Y A L D Y N E Y F R S V G Y K P D F V G F E I P D K F V V G Y A L D Y N E Y F R S V G Y R P D F V G F E I P D K F V V G Y A L D Y N E Y F	· · ·	240		S	⋖			
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FIG.3C

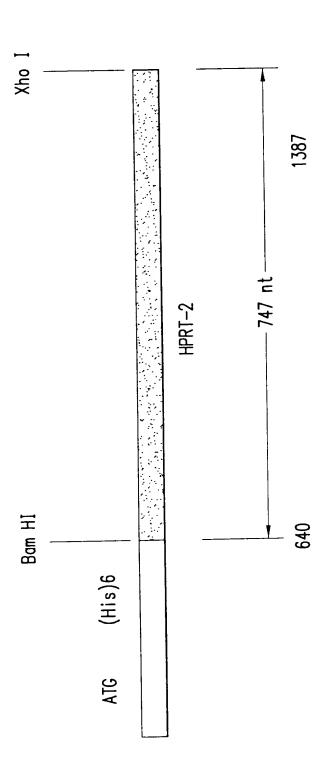


FIG.4

